

Nov 23 14:57

us-08-913-430-1.mli

1

GenCore version 4.5  
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OM nucleic - nucleic search, using ew model

Run on: November 19, 1999, 23:20:33 ; Search time 29.19 Seconds  
(without alignments)  
6455.963 Million cell updates/sec

Title: US-08-913-430-1

Perfect score: 1782

Sequence: 1 atgaaaaaatgccactata.....ttgataactcgtatgctctaa 1782

Scoring table: IDENTITY\_NUC

Searched: 195662 seqs, 52875861 residues

Database :

- Issued\_Patents\_NA.\*
- 1: /cgn2\_6/ptodata/1/ina/5a\_COMB.seq.\*
  - 2: /cgn2\_6/ptodata/1/ina/5b\_COMB.seq.\*
  - 3: /cgn2\_6/ptodata/1/ina/5c\_COMB.seq.\*
  - 4: /cgn2\_6/ptodata/1/ina/5d\_COMB.seq.\*
  - 5: /cgn2\_6/ptodata/1/ina/PCTUS9.COMB.seq.\*
  - 6: /cgn2\_6/ptodata/1/ina/backfile1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Match	Length	DB	ID	Description
1	77.4	4.3	1511	1	US-07-991-867B-8	Sequence 8, Appl
2	77.4	4.3	1511	2	US-08-107-755A-8	Sequence 8, Appl
3	77.4	4.3	1511	4	US-08-544-332-8	Sequence 8, Appl
4	75	4.2	5852	1	US-07-867-106-2	Sequence 2, Appl
5	74.4	4.2	8920	3	US-08-446-855A-1	Sequence 1, Appl
6	72.6	4.1	6243	4	US-09-056-075-1	Sequence 1, Appl
7	71	4.0	731	2	US-08-451-405A-2	Sequence 2, Appl
8	69.2	3.9	1939	1	US-07-715-751B-2	Sequence 2, Appl
9	68.2	3.8	731	2	US-08-451-405A-2	Sequence 1, Appl
10	66	3.7	660	1	US-07-991-867B-32	Sequence 32, Appl
11	66	3.7	660	2	US-08-107-755A-32	Sequence 32, Appl
12	66	3.7	660	4	US-08-544-332-32	Sequence 32, Appl
13	64.6	3.6	6243	4	US-09-056-075-1	Sequence 1, Appl
14	64.4	3.6	5852	1	US-07-867-106-2	Sequence 2, Appl
15	63	3.5	8457	1	US-07-991-867B-1	Sequence 1, Appl
16	63	3.5	1395	1	US-07-991-867B-25	Sequence 25, Appl
17	63	3.5	6768	2	US-08-107-755A-1	Sequence 1, Appl
18	63	3.5	1395	2	US-08-107-755A-25	Sequence 25, Appl
19	63	3.5	8457	4	US-08-544-332-1	Sequence 1, Appl
20	63	3.5	1395	4	US-08-544-332-25	Sequence 25, Appl
21	62.4	3.5	1511	1	US-07-991-867B-8	Sequence 8, Appl
22	62.4	3.5	240	1	US-08-628-417-6	Sequence 6, Appl
23	62.4	3.5	1511	2	US-08-107-755A-8	Sequence 8, Appl
24	62.4	3.5	1511	4	US-08-544-332-8	Sequence 8, Appl
25	61.8	3.5	1939	1	US-07-715-751B-2	Sequence 2, Appl
26	61.8	3.5	4098	4	US-08-605-106-4	Sequence 4, Appl
27	61.6	3.5	660	1	US-07-991-867B-32	Sequence 32, Appl
28	61.6	3.5	660	2	US-08-107-755A-32	Sequence 32, Appl

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29	61.6	3.5	660	4	US-08-544-332-32	Sequence 32, Appl
30	60.8	3.4	8920	3	US-08-446-855A-1	Sequence 1, Appl
31	60.6	3.4	1079	1	US-07-781-355-1	Sequence 1, Appl
32	60.4	3.4	3138	1	US-07-867-106-4	Sequence 4, Appl
33	60.4	3.4	3618	1	US-07-872-678A-36	Sequence 36, Appl
34	60.2	3.4	2447	4	US-09-014-969-14	Sequence 14, Appl
35	60	3.4	1422	3	US-08-319-704-5	Sequence 5, Appl
36	59	3.3	1241	1	US-07-593-657-6	Sequence 16, Appl
37	58.6	3.3	8700	3	US-08-392-625-16	Sequence 16, Appl
38	58.6	3.3	8700	3	US-08-466-961A-16	Sequence 16, Appl
39	58.6	3.3	8700	4	US-08-645-193B-18	Sequence 18, Appl
40	58.4	3.3	8457	1	US-07-991-867B-1	Sequence 1, Appl
41	58.4	3.3	6768	2	US-08-107-755A-1	Sequence 1, Appl
42	58.4	3.3	8457	4	US-08-544-332-1	Sequence 1, Appl
43	58.2	3.3	4970	3	US-08-764-100-14	Sequence 14, Appl
44	58.2	3.3	4970	3	US-08-764-100-20	Sequence 20, Appl
45	58	3.3	2334	2	US-08-062-632-4	Sequence 4, Appl

#### ALIGNMENTS

RESULT 1  
US-07-991-867B-8/c  
Sequence 8, Application US/07991867B  
Patent No. 5476781

GENERAL INFORMATION:  
APPLICANT: Moyer, Richard W.  
APPLICANT: Hall, Richard L.  
APPLICANT: Gruidl, Michael E.  
TITLE OF INVENTION: No. 5476781el Entomopoxvirus Expression System  
NUMBER OF SEQUENCES: 66  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: David R. Salivanchik  
STREET: 2421 N.W. 41st Street, Suite A-1  
CITY: Gainesville  
STATE: FL  
COUNTRY: USA  
ZIP: 32606  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/07/991,867B  
FILING DATE: 12-DEC-1992  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: WO 92/14818  
FILING DATE: 12-FEB-1992  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/827,685  
FILING DATE: 30-JAN-1992  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/657,584  
FILING DATE: 19-FEB-1991  
ATTORNEY/AGENT INFORMATION:  
NAME: Salivanchik, David R.  
REGISTRATION NUMBER: 31,794  
REFERENCE/DOCKET NUMBER: UFI14.C3  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 904-375-8100  
TELEFAX: 904-372-5800

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us-08-913-430-1.mli

3

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; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
;   LENGTH: 1511 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: double
;   TOPOLOGY: unknown
;   MOLECULE TYPE: DNA (genomic)
;   ORIGINAL SOURCE:
;     ORGANISM: Amsacta moorei entemopoxvirus
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: complement (18..218)
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: complement (234..782)
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: 852..1511
;   US-07-991-867B-8
;
Query Match      4.3%; Score 77.4; DB 1; Length 1511;
Best Local Similarity 49.8%; Pred. No. 9.5e-06;
Matches 341; Conservative 0; Mismatches 326; Indels 18; Gaps 5;
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Qy 152 aaaaataaagttctatctatctctctctctctctctctctctctctctctct 211
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Db 1248 ATATTAATTAATTTTAAATTTAAATTTAAATTTAAATTTAAATTTAAATTT 1189
Qy 212 agtattatcaagtcggtatttttctatctctctctctctctctctctctct 271
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Db 1188 ATGATTCACATTTTAAATTTCTATGAGTGTGCTATGATTTATTTATTTGCT 1129
Qy 272 ttccataatcctaataatttctatctctctctctctctctctctctctctct 330
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Qy 331 ttatagatttttttttttttttttttttttttttttttttttttttttttt 390
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Db 1068 TAAATTAATTTTAAATTTGAAATATATTAATTTTAAATTTTCTGAAATAT 1009
Qy 391 tttaataattagtaaaaaatacaaatat-----tttcctattcagaataatc 446
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Db 888 TATTTTATTAATTAATATATGCAATTTGAAATTCATTTGATTAATTTTATTT 829
Qy 561 agctattatgcaactcgtctgcatcaattatgcatctgtgcaagcagtgtaga 620
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Db 828 ATTGATTT-----AATTTTATTTTGAATAAATATATCAATTAATTAATTA 774
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Db 773 GAATTAATTAATTTGCTCTATGTTTCTGCAAAACAAAGAAATGATGCAAAATTA 714
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us-08-913-430-1.mli

4

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Qy 679 graagtaatgattcattcgatgaatagcaacgatccgagtaactcctcgatgaatt 738
    || || || || || || || || || || || || || || || || || || ||
Db 713 AGATATATTTTAAATTTCTAATCAAAATGCTAATTTAACTCATTAATATATGATTT 654
Qy 739 gcccaaaaagattatttctctatg 763
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Db 653 ATTAATAAATAATTAATTAATCATG 629
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RESULT 2
US-08-107-755A-8/c
; Sequence 8, Application US/08107755A
; Patent No. 5721352
; GENERAL INFORMATION:
; APPLICANT: Moyer, Richard W.
; APPLICANT: Hall, Richard L.
; APPLICANT: Gruidl, Michael E.
; TITLE OF INVENTION: No. 5721352el Entomopoxvirus Expression System
; NUMBER OF SEQUENCES: 40
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: David R. Saliwanchik
; STREET: 2421 N.W. 41st Street, Suite A-1
; CITY: Gainesville
; STATE: Florida
; COUNTRY: U.S.A.
; ZIP: 32606
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/107, 755A
; FILING DATE: 19-AUG-1993
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/827, 658
; FILING DATE: 30-JAN-1992
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/657, 584
; FILING DATE: 19-FEB-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Saliwanchik, David R.
; REGISTRATION NUMBER: 31, 794
; REFERENCE/DOCKET NUMBER: UF114.C2
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (904) 375-8100
; TELEFAX: (904) 372-5800
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
;   LENGTH: 1511 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: double
;   TOPOLOGY: unknown
;   MOLECULE TYPE: DNA (genomic)
;   ORIGINAL SOURCE:
;     ORGANISM: Amsacta moorei entemopoxvirus
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: complement (18..218)
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: complement (234..782)
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Query Match	4.3%;	Score 77.4;	DB 2;	Length 1511;
Best Local Similarity	49.8%;	Pred. No. 9.5e-06;		
Matches 341;	Conservative	0;	Mismatches 326;	Indels 18;
				Gaps 5;

Db 653 ATAAATAAAATATAATAATCATG 629

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RESULT      3
US-08-544-332-8/c
; Sequence 8, Application US/08544332
; Patent No. 5935777
; GENERAL INFORMATION:

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ADDRESSEE: Gerard H. Bencen  
STREET: 2421 N.W. 41st Street, Suite A-1  
CITY: Gainesville  
STATE: FL

COMPUTER READABLE FORM:

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; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
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; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25

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; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/544,332

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; FILING DATE:
; CLASSIFICATION: 435
;

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; PRIOR APPLICATION DATA:  
 ; APPLICATION NUMBER: US 07/991,867

; FILING DATE: 07-DEC-1992  
 ; PRIOR APPLICATION DATA:  
 ;

APPLICATION NUMBER: US 0  
FILING DATE: 19-AUG-1993

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: WO 92/14818

FILING DATE: 12-FEB-1992  
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/827,685  
FILING DATE: 30-JAN-1992

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/657,584

FILING DATE: 19-FEB-1991  
ATTORNEY/AGENT INFORMATION:

NAME: Bencen, Gerard H.  
REGISTRATION NUMBER: 35.746

REFERENCE/DOCKET NUMBER:  
TELECOMMUNICATION INFORMATION

TELEPHONE: 904-375-8100  
TELEFAX: 904-372-5800

1 TELEFAX. 504-572-5000  
2  
3 INFORMATION FOR SEQ ID NO: 8  
4  
5 SEQUENCE CHARACTERISTICS:

SEQUENCE CHARACTERISTICS:  
; LENGTH: 1511 base pairs  
; TYPE: nucleic acid

STRANDEDNESS: double  
; nucleic acid  
; topology: unbranched

MOLECULE TYPE: DNA (genomic)  
; ORIGINAL SUBJECT:

ORIGINAL SOURCE:  
; ORGANISM: Amsacta moorei entomopoxvirus  
; ;  
END.

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; FEATURE:
;
; NAME/KEY: CDS
; LOCATION: complement (19

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LOCATION: complement (10...210)
FEATURE:
NAME/KEY: CDC

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NAME/KEY: CDS
LOCATION: complement (234..782)
END
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:	NAME/KEY:	CDS
:	LOCATION:	DEC 1971

LOCATION: 852..1311  
US-08-544-332-8









5

RESULT 10  
US-07-991-867B-32/c  
; Sequence 32, Application US/07991867B  
; Patent No. 5476781  
; GENERAL INFORMATION:  
; APPLICANT: Moyer, Richard W.  
; APPLICANT: Hall, Richard L.  
; APPLICANT: Grudl, Michael E.  
; TITLE OF INVENTION: No. 5476781e1 Entomopoxvirus Expression System  
; NUMBER OF SEQUENCES: 66  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: David R. Saliwanchik

	Query Match	3.78;	Score 66;	DB 1;	Length 660;	
	Best Local Similarity	51.29;	Pred. No. 0.00076;			
	Matches 205;	Conservative	0;	Mismatches 190;	Indels	Gaps
Qy	92	tagtacattaaaagtcagaacaaagtattcaatgtaactgtgaaccttagcgccatccttaagaa	151			
Db	457	TATTTAACACTAATCTAACTAATAGAAAAACATTATTAAGTTAACCAATTATTTTGG	398			
Qy	152	aaatataaagtttcatcctatctttttttaaocgaataccacaaggcataaatcttgctc	211			
Db	397	ATATTATTATTTTTTTAAATTTCATAAATTATTATTAATAATTATAGCATTTATATTAC	338			
Qy	212	agatttaacsaagtcgggatatttttcatattcttctaactaaatattatgaattgcgat	271			
Db	337	ATGATTGCACATTTAAAATAATTCGTATGCAATCGTGTCGTATGAATTTACTGTATTCGTAT	278			
Qy	272	tcttcataatcctaataattcacattttt- <del>cata</del> ccaattttttaaactttttaaactccttcaat	330			
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US-08-913430-1.mh

6

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? FILING DATE: 30-JAN-1992
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US 07/657,584
? FILING DATE: 19-FEB-1991
? ATTORNEY/AGENT INFORMATION:
? NAME: Bencken, Gerard H.
? REGISTRATION NUMBER: 35,746
? REFERENCE/DOCKET NUMBER: UFI14,C4
? TELECOMMUNICATION INFORMATION:
? TELEPHONE: 904-375-8100
? TELEFAX: 904-372-5800
? INFORMATION FOR SEQ ID NO: 32:
? SEQUENCE CHARACTERISTICS:
? LENGTH: 660 base pairs
? TYPE: nucleic acid
? STRANDEDNESS: double
? TOPOLOGY: unknown
? MOLECULE TYPE: DNA (genomic)
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US-08-544-332-32

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Nov 23 14:57

US-68-913-430-1 PH

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1  NUMBER OF SEQUENCES: 2
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3  CORRESPONDENCE ADDRESS:
4  ADDRESSEE: Quarles & Brady
5  STREET: 1 South Pinckney Street
6  CITY: Madison
7  STATE: WI
8  COUNTRY: US
9  ZIP: 53701-2113
10
11  COMPUTER READABLE FORM:
12  MEDIUM TYPE: Floppy disk
13  COMPUTER: IBM PC compatible
14  OPERATING SYSTEM: PC-DOS/MS-DOS
15  SOFTWARE: PatentIn Release #1.0, Version #1.30
16  CURRENT APPLICATION DATA:
17  APPLICATION NUMBER: US/09/056,075
18  FILING DATE:
19
20  CLASSIFICATION:
21  ATTORNEY/AGENT INFORMATION:
22  NAME: Seay, Nicholas J
23  REGISTRATION NUMBER: 27386
24  REFERENCE/DOCKET NUMBER: 960296,95238
25  TELECOMMUNICATION INFORMATION:
26  TELEPHONE: 608-251-5000
27  TELEFAX: 608-251-9166
28  INFORMATION FOR SEQ ID NO: 1:
29
30  SEQUENCE CHARACTERISTICS:
31  LENGTH: 6243 base pairs
32  TYPE: nucleic acid
33  STRANDEDNESS: double
34  TOPOLOGY: Linear
35  MOLECULE TYPE: DNA (genomic)
36  FEATURE:
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38  NAME/KEY: misc.feature
39  LOCATION: 3770..4013
40  OTHER INFORMATION: /note="RP4 origin of DNA transfer (orit) from
41  OTHER INFORMATION: plasmid RP4"
42
43  US-09-056-075-1

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Nov 23 14:57

us-08-913-430-1.mi

23

:/ PRIOR APPLICATION DATA:  
:/ APPLICATION NUMBER: US 07/827, 685  
:/ FILING DATE: 30-JAN-1992  
:/ PRIOR APPLICATION DATA:  
:/ APPLICATION NUMBER: US 07/657, 584  
:/ FILING DATE: 19-FEB-1991  
:/ ATTORNEY/AGENT INFORMATION:  
:/ NAME: Saliwanchik, David R.  
:/ REGISTRATION NUMBER: 31,794  
:/ REFERENCE/DOCKET NUMBER: UFI14.C3  
:/ TELECOMMUNICATION INFORMATION:  
:/ TELEPHONE: 904-375-8100  
:/ TELEFAX: 904-372-5800  
:/ INFORMATION FOR SEQ ID NO: 1:  
:/ SEQUENCE CHARACTERISTICS:  
:/ LENGTH: 8457 base pairs  
:/ TYPE: nucleic acid  
:/ STRANDEDNESS: double  
:/ TOPOLOGY: unknown  
:/ MOLECULE TYPE: DNA (genomic)  
:/ ORIGINAL SOURCE:  
:/ ORGANISM: Amsacta moorei entomopoxvirus  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: complement (65..1459)  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: 1474..2151  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: complement (2239..2475)  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: 2502..2987  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: 3080..6091  
:/ FEATURE:  
:/ NAME/KEY: CDS  
:/ LOCATION: complement (6277..6768)  
:/ US-07-991-867B-1

Query Match 3.5%; Score 63; DB 1; Length 8457;  
Best Local Similarity 48.2%; Pred. No. 0.0031;  
Matches 206; Conservative 0; Mismatches 220; Indels 1; Gaps 1;  
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Db 623 TAGTACTGCTTAATCCAGAACTTATAAACAATAAATAATATTTATTGATC 564  
Qy 179 taatgaatccaccagcataaactttgtcagttatttcaagtcogttatttca 238  
Db 563 TGATATGACTTATAAAGCATAA-CAGTACAAATGTAAGTCTTAATACATAATGCTT 505  
Qy 239 ttatttcacaaatattattgaatttgatttcataatcctaaatttaccttt 298  
Db 504 ATTATACATTGATGTAATCAATTAATTAATGATATTTAAATGTAATTAAT 445  
Qy 299 ttataacaattttaaatctttaaattatagatatttttattttttagctca 358  
Db 444 ACTTTTAAATACATTGTAATTAATGATGATGATGATGATGATGATGATGAT 385  
Qy 359 aattataaatatcttgaatttattgaatttataatttagtactcaaaaatacaa 418

Nov 23 14:57

us-08-913-430-1.mi

24

Db 384 TATTTTATATATATATAGTTTATTTCAACATTAATCTMAATTTCAATTTAAATTAAT 325  
Qy 419 atatttttctatctcaagaaatcattttttaaaaaattgattttatagata 478  
Db 324 ATTATTCATGAGTTTATTGCTGATTTATGCTAGTACTTATAGAGTGCATTAATTA 265  
Qy 479 attgttgtaaatgaatlaacttgatttgaaggaacaaatgaaaaatgctta 538  
Db 264 TTCTAATATGATATTTAATAGTATTCAGCAATTTAAATAATCATTTAATA 205  
Qy 539 gaaaaa 545  
Db 204 TAAAAA 198

Search completed: November 20, 1999, 00:02:31  
Job time: 2518 sec

Nov 23 14:57

us-08-913-430-1.mmp

1

GenCore version 4.5  
Copyright (c) 1993 - 1998 CompuGen Ltd.

OM nucleic - nucleic search, using sw model

Run on: November 19, 1999, 23:06:31 ; Search time 450.03 Seconds  
(without alignments)  
12182.104 Million cell updates/sec

Title: US-08-913-430-1

Perfect score: 1782

Sequence: 1 atgaaaaaatgccactata.....ttgataatcctgatgcctaa 1782

Scoring table: IDENTITY\_NUC

Searched: 4116993 seqs, 1538246968 residues

Database :

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1: /cgn2\_6/ptodata/1/pna/PCTUS9\_COMB.seq:\*  
2: /cgn2\_6/ptodata/1/pna/US07\_COMB.seq:\*  
3: /cgn2\_6/ptodata/1/pna/US080\_COMB.seq:\*  
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2

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result		Query		DB	ID	Description
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1	1782	100.0	1782	26	US-08-913-430-1	Sequence 1, Appli
2	1245.6	69.9	1260	7	US-08-183-714-2	Sequence 2, Appli
3	97.4	5.5	448	80	US-60-141-136-611	Sequence 611, App
4	95.4	5.4	17034	80	US-60-144-351-1275	Sequence 1275, Ap
5	93.2	5.2	501	80	US-60-141-134-4006	Sequence 4006, A
6	92.6	5.2	470	81	US-60-145-485-8111	Sequence 8111, Ap
7	92.2	5.2	510	80	US-60-144-084-39263	Sequence 39263, A
8	92	5.2	837	33	US-08-998-416-288	Sequence 288, App
9	91.8	5.2	25821	88	US-60-161-932-409	Sequence 409, App
10	91.6	5.1	1431	77	US-60-126-265-4175	Sequence 4175, Ap

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c 13	91.4	5.1	961	77	US-60-126-265-3039	Sequence 3039, A
c 14	90.8	5.1	948	52	US-09-404-520-13069	Sequence 13069, A
c 15	89.8	5.0	1074	81	US-60-146-055-46	Sequence 46, Appl
c 16	89.4	5.0	507	80	US-60-141-134-1097	Sequence 1097, Ap
c 17	89.2	5.0	529	77	US-60-128-440-1802	Sequence 1802, Ap
c 18	89	5.0	445	77	US-60-128-440-218	Sequence 218, App
c 19	89	5.0	497	81	US-60-145-485-5711	Sequence 5711, Ap
c 20	89	5.0	430	81	US-60-145-485-7216	Sequence 7216, Ap
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c 22	88.4	5.0	433	81	US-60-145-485-8223	Sequence 8223, Ap
c 23	88.4	5.0	447	81	US-60-145-485-9030	Sequence 9030, Ap
c 24	87.6	4.9	908	77	US-60-126-265-3038	Sequence 3038, Ap
c 25	87.4	4.9	400	80	US-60-141-134-1072	Sequence 1072, Ap
c 26	87.4	4.9	1748	80	US-60-144-351-2763	Sequence 2763, Ap
c 27	86.8	4.9	17034	80	US-60-144-351-1275	Sequence 1275, Ap
c 28	86.6	4.8	570	80	US-60-141-134-2612	Sequence 2612, Ap
c 29	86.4	4.8	458	80	US-60-141-136-716	Sequence 716, App
c 30	86.4	4.8	667	52	US-09-404-520-13658	Sequence 13658, A
c 31	86.2	4.8	433	80	US-60-141-134-3384	Sequence 3384, Ap
c 32	85.4	4.8	509	81	US-60-145-146-991	Sequence 991, App
c 33	85.4	4.8	636	33	US-08-998-416-1137	Sequence 1137, Ap
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c 35	84.6	4.7	19124	15	US-08-487-826-13	Sequence 13, Appl
c 36	84	4.7	19124	15	US-08-487-826-13	Sequence 13, Appl
c 37	84	4.7	19124	15	US-08-487-826-13	Sequence 13, Appl
c 38	84	4.7	19124	15	US-08-487-826-13	Sequence 13, Appl
c 39	83.8	4.7	805	77	US-60-126-265-127	Sequence 127, App
c 40	83.4	4.7	1174	52	US-09-404-520-13397	Sequence 13397, A
c 41	83	4.7	463	81	US-60-145-147-1007	Sequence 1007, Ap
c 42	83	4.7	378	81	US-60-145-485-6622	Sequence 6622, Ap
c 43	82.8	4.6	9734	14	US-08-430-908-20	Sequence 20, Appl
c 44	82.8	4.6	9741	17	US-08-639-368-43	Sequence 43, Appl
c 45	82.8	4.6	9741	17	US-08-639-368-43	Sequence 43, Appl

ALIGNMENTS

RESULT 1  
US-08-913-430-1  
Sequence 1, Application US/08913430B

GENERAL INFORMATION:  
APPLICANT: Walker, John

APPLICANT: Lee, Rogan

APPLICANT: Dougherty, Stephen W.

FILE REFERENCE: D-011415-0

CURRENT APPLICATION NUMBER: PCT/US96/00149

EARLIER FILING DATE: 1996-03-15

EARLIER APPLICATION NUMBER: PN 1789

NUMBER OF SEQ ID NOS: 18

SOFTWARE: PatentIn Ver. 2.0 - beta

SEQ ID NO 1

TYPE: DNA

ORGANISM: Mycoplasma hyopneumoniae

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RESULT 2

US-08-183-774-2

; Sequence 2, Application US/08183774

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/ GENERAL INFORMATION:
/ APPLICANT: SETO, Yasuhiro
/ APPLICANT: FUTO, Satoshi
/ APPLICANT: MITSUSE, Shizuo
/ APPLICANT: MATSUI, Kanako
/ APPLICANT: TSUNA, Mika
/ TITLE OF INVENTION: DNA's Encoding Surface Antigen of
/ TITLE OF INVENTION: Mycoplasma hyopneumoniae, DNA Fragments for Primer,
/ TITLE OF INVENTION: Recombinant Antigenic Peptides and Diagnostic Method of
/ NUMBER OF SEQUENCES: 18
/ CORRESPONDENCE ADDRESS:
/ ADDRESSEE: OBLON, SPIVAK, MCLELLAND, MAIER & NEUSTADT,
/ ADDRESSEE: P.C.
/ STREET: 1755 Jefferson Davis Highway, Fourth Floor
/ CITY: Arlington
/ STATE: Virginia
/ ZIP: 22202
/ COMPUTER READABLE FORM:
/ MEDIUM TYPE: Floppy disk
/ COMPUTER: IBM PC compatible
/ OPERATING SYSTEM: PC-DOS/MS-DOS
/ SOFTWARE: PatentIn Release #1.0, Version #1.25
/ CURRENT APPLICATION DATA:
/ APPLICATION NUMBER: US/08/183,774
/ FILING DATE:
/ CLASSIFICATION: 435
/ PRIOR APPLICATION DATA:
/ APPLICATION NUMBER: US 07/747,015
/ FILING DATE: 19-AUG-1991
/ ATTORNEY/AGENT INFORMATION:
/ NAME: Oblon, Norman F.
/ REGISTRATION NUMBER: 24, 618
/ REFERENCE/DOCKET NUMBER: 4183-001-0
/ TELECOMMUNICATION INFORMATION:
/ TELEPHONE: (703)521-4500
/ TELEFAX: (703)486-2347
/ TELEX: 248855 OPAT UR
/ INFORMATION FOR SEQ ID NO: 2:
/ SEQUENCE CHARACTERISTICS:
/ LENGTH: 1260 base pairs
/ TYPE: nucleic acid
/ STRANDEDNESS: single
/ TOPOLOGY: linear
/ MOLECULE TYPE: DNA (genomic)
/ ORGANISM: Mycoplasma hyopneumoniae
/ STRAIN: J
/ US-08-183-774-2

Query Match 69.9%; Score 1245.6; DB 7; Length 1260;
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Matches 1251; Conservative 0; Mismatches 9; Indels 0; Gaps 0;

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Qy 583 gcatcaattatgcatgttgcgcaggttctggaagacagacatcaggttaactct 642
    |||||||
Db 61 GCATCAATATTGCAATTTTGTGCACAGGTTGCGACAGCAAGAAATCAGCTTCCTCA 120

Qy 643 gatctaaccacgaagcgagacgctaacaacataaagtaagtaagtaagtaagtaagta 702
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[illegible]

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Db	1141	CCAAATATTTCATTTAAATATGATTAATCAACATATPAAGCGAAGTAAATAATTAAT	1200
Qy	1723	acaatttgaagtcgcgatattgttaaaaagcraatgltgataatccgatgactaa	1782
Db	1201	ACAATTTTATGTAATCGCATATTTGTAACAAAGCTATPAAGTTGATATCTGTATGCCCTA	1260

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RESULT 3
US-60-141-136-611
; Sequence 611, Application US/60141136
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S.
; APPLICANT: Ia Rosa, Thomas J.
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(15709)A
; CURRENT FILING DATE: 1999-06-28
; NUMBER OF SEQ ID NOS: 4228
; SEQ ID NO 611
; LENGTH: 448
; TYPE: DNA
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB3108-008-Q1-K1-H6
US-60-141-136-611

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Oy	228 tatttttcaatctactcctaataatatttggaattgcatlittccaalctcaaat	288				
Dd	79 tttaatttttttttttttttttttttttttttttttccttataggittaataaaaaaaataa	138				
Oy	289 ttacaccttttttlataacaattttaaaaactcactcttaattatagratittttlat	348				
Dd	133 aaatraattttttttaaaaaaaaaaaaaaaaaaaaaaaaaaatagtcgycgcgt	198				
Oy	349 tttagtgtcaaattraatatraccttgatattgatggattttttaaattagtracta	408				
Dd	199 ttaaaataaaaaaataaataaaccttcottttaaataaatalttttttttttaaa	258				
Oy	409 aaaaatacaatatlttttccctatctcaagaanaatcatlltttaaaaaaatgatl	468				
Dd	259 taasaaaaaataatttaccitctttaaaaaaataataaaaaaaataatta	318				
Oy	465 ttatagtataattgttqgtataatgnaatcaactgabtgygaagygaacaatgaa	528				
Dd	319 taataataaataatattatctataaaaaaatattttcttatgtgaattataaaaaaa	378				
Oy	529 aaaatgctgaagaaaatcttc	551				
Dd	379 aaaaataatcaaatltttt	401				

RESULT 4  
US-60-144-351-1275



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```
; Sequence 1275, Application US/60144351
; GENERAL INFORMATION:
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Kovalic, David K.
; APPLICANT: Liu, Jindong
; TITLE OF INVENTION: Plant Genome Sequence And Uses Thereof
; FILE REFERENCE: 38-21(51237)A
; CURRENT APPLICATION NUMBER: US/60/144,351
; CURRENT FILING DATE: 1999-07-20
; NUMBER OF SEQ ID NOS: 5763
; SEQ ID NO 1275
; LENGTH: 17034
; TYPE: DNA
; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: unsure at all n locations
US-60-144-351-1275
```

```
Query Match          5.4%; Score 95.4; DB 80; Length 17034;
Best Local Similarity 46.6%; Pred. No. 8.7e-08;
Matches 244; Conservative 0; Mismatches 268; Indels 12; Gaps 1;

Qy 81 ttttaagaattgacatcaataaagtagcaacaagtatcaatgtaaacattagcgca 140
    |||||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 80 ttttttttaaaaaataaaaaaannatttttaaaaaatttttttttttttttttttt 139

Qy 141 acccttaagaanaatttaaaagttatctatcttttttaacgaaatccaccgagcat 200
    |  ||  |||||  ||  |||  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 140 ammmmaaaaaaataaataatttttttttttttttttttttttttttttttttttttt 199

Qy 201 aaactcttgacgtattatcaagtcggtatcttttttttttttttttttttttttttt 260
    |||||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 200 aaatnaammmmaaaatnaaaaaatnaaaatttttaatatattttaaanaaatTTTTTT 259

Qy 261 tgaattgcatttccatcaatctaaattttacatttttttttttttttttttttttttt 313
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 260 ttattgtttataaaaaaatacaaaaaaanntttttttttttttttttttttttttttt 319

Qy 313 -----taaaaatctcttaattatagatttttttttttttttttttttttttttttttt 368
    |||||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 320 nmmmaaaaaaataaaaaaannmmatttttttttttttttttttttttttttttttttt 379

Qy 369 ttatctgaattttattgaatttttataatttagtactaaaaatcaaatatcttttc 428
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 380 aaaaaaaatannnttttttttttttttttttttttttttttttttttttttttttttt 439

Qy 429 ctatctcagaanaatcatttttttttttttttttttttttttttttttttttttttttt 488
    ||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 440 ttaattttaaatataaaaaaataaannltttaaannatatammnnmttttt 499

Qy 489 ataattgaatnaacttgatttgaagggacaacaatgaaaaaactgtagaanaaatt 548
    |||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 500 atataataatataatataatataatataanaaatatataatataataaaaaa 559

Qy 549 ctgtatcatcagctattatgaacttgctgtcatcaat 592
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 560 aaaaamatatataatataatatttttntntnmma 603
```

```
RESULT 5
US-60-141-134-4006
; Sequence 4006, Application US/60141134
; GENERAL INFORMATION:
```

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```
; APPLICANT: Bewley, Derek J.
; APPLICANT: Deikman, Jill
; APPLICANT: Thompson, Michael D.
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; FILE REFERENCE: 38-21(15137)A
; CURRENT APPLICATION NUMBER: US/60/141,134
; CURRENT FILING DATE: 1999-06-28
; NUMBER OF SEQ ID NOS: 4434
; SEQ ID NO 4006
; LENGTH: 501
; TYPE: DNA
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: LIB3094-095-Q1-K1-H9
US-60-141-134-4006
```

```
Query Match          5.2%; Score 93.2; DB 80; Length 501;
Best Local Similarity 51.9%; Pred. No. 1e-07;
Matches 258; Conservative 0; Mismatches 234; Indels 5; Gaps 2;

Qy 77 agaatttttaagaattgacatcaataaagtagcaacaagtatcaatgtaaacattag 136
    |  |||||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 6  aaatttttttaaaaaaataaaaaaataaataattgttttttttttttttttttttttttt 65

Qy 137 cgaactcttaagaanaatttaaaagttatctatcttttttttttttttttttttttttttt 196
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 66 tdtgtggcccaaaaaaataaataattttatataatataattctatcatcaataa 125

Qy 197 gcaaatctcttgacgtattatcaagtcggtatcttttttttttttttttttttttttttt 256
    |||  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 126 ttatagaaatctactaatat-aaatatttaataatataatatttaaatat 184

Qy 257 tatgttgattgcatttccatcaatctaaattttacatttttttttttttttttttttttt 312
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 185 tctttttataaaaatatttttttttttttttttttttttttttttttttttttttttttt 244

Qy 313 taaaatactctttaaattatagatttttttttttttttttttttttttttttttttttttt 372
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 245 tgaataaatctatttaatttttttttttttttttttttttttttttttttttttttttt 304

Qy 373 ctgaattttattgaatttttataatttagtactaaaaatcaaatatttttctat 432
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 305 aataaataatatttaattatatttttttttttttttttttttttttttttttttttttttt 364

Qy 433 tcttaagaanaatcatttttaaaaaaattgattttatagatataattgtttgataa 492
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 365 taatatataataataatatttaatttaataataataatatttaatttttttttttttt 424

Qy 493 ttgaatttaattgatttgaagggacaacaatgaaaaaactgtagaanaaattctg 552
    ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||  ||
Db 425 ttaaaaaaatataatataataaataacattttatatacaaatattttatataaatttaatt 484

Qy 553 tattatcagctattta 569
    |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
Db 485 atatactattattaa 501
```

```
RESULT 6
US-60-145-485-8111/c
; Sequence 8111, Application US/60145485
; GENERAL INFORMATION:
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APPLICANT: Steiner, Sabine  
 APPLICANT: Mohr, Christine  
 APPLICANT: Wendland, Jurgen  
 APPLICANT: Knechtel, Philipp  
 APPLICANT: Reibschung, Corinne  
 TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSYPII  
 TITLE OF INVENTION: AND USES THEREOF  
 NUMBER OF SEQUENCES: 1152  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Novartis Corporation  
 STREET: 3054 Cornwallis Road  
 CITY: Research Triangle Park  
 STATE: North Carolina  
 COUNTRY: USA  
 ZIP: 27709  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: PatentIn Release #1.0, Version #1.30  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/98,416  
 FILING DATE: 24-DEC-1997  
 CLASSIFICATION: 435  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: CH 0016/97  
 FILING DATE: 31-DEC-1996  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Meigs, J. Timothy  
 REGISTRATION NUMBER: 38,241  
 REFERENCE/DOCKET NUMBER: PF/5-30306/A/GC1976  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 919-541-8587  
 TELEFAX: 919-541-8689  
 INFORMATION FOR SEQ ID NO: 288:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 837 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: Linear  
 MOLECULE TYPE: DNA (genomic)  
 ORIGINAL SOURCE:  
 ORGANISM: PAB1241RP

Query Match	5.2%;	Score 92;	DB 33;	Length 837;
Best Local Similarity	48.9%;	Pred. No. 1.8e-07;		
Matches 305; Conservative	0;	Mismatches 315;	Indels 4;	Gaps 21

[illegible]

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Db	201	ATATGAAATACATTTTGGCTAGTCGAATTTTTAAATTGATTATTAATAAATATTATGTAT	260
Oy	318	attactcttaattatagatatttttatttttagtcctaataatcataatcgtga	377
Db	261	ATTATATTTTCTTTTAATAATATTTAAATAGATTA---TGAAATATTAATATATATTT	317
Oy	378	atttatitgaaatttttaaatattagtaacataaataatltttccattcaa	437
Db	318	ATTATATGTTATTAATAAATAATATATTTATTTATTAAGAATTAATTTATTAATAT	377
Oy	438	gaaaaatcatttttttaaaaaaatigtatttttaagtataatttggttgataattgaa	497
Db	378	TGTAAATATTTATTTATTTATTAATAATCTAATTTTAAATATTTGTCATTTATTAATTA	437
Oy	498	ttaacttgtttgaaggacaacaataagaanaaacctcagaanaaacctctgcatac	557
Db	438	TTTATCTTTTATTAACAATTTATTTAAATTAATTTTAACTTAATTTCTATTAATTA	497
Oy	558	atcacgattatccaactcgcttgcatacaattatgcatttggcagcaggttg	617
Db	498	ATTATTAATTAATTAATAATTAATTAATTCATTTATTTATTTATTTATTAATTAATTT	557
Oy	618	acagacagaaatcaggticaactcttgatcttaaaccagaacgcagacgtcaaacataa	677
Db	558	AATTAATTAATTAATTAATTAATTAATTCATTTATTTAAATTAATAATTAATTAAGAATGT	617
Oy	678	agtaagaatcaggtcattcgaat	701
Db	618	ACTTAAATAATCTATTAAGAAT	641

```

RESULT          9
; Sequence 409, Application US/60161932
; GENERAL INFORMATION:
; APPLICANT: Venter, J. Craig
; TITLE OF INVENTION: The Drosophila Genome, Primary Nucleic
; TITLE OF INVENTION: Acid Sequences, Systems Containn the Nucleic Acid Sequences
; TITLE OF INVENTION: and Uses Thereof
; FILE REFERENCE: CL000122
; CURRENT APPLICATION NUMBER: US/60/161,932
; CURRENT FILING DATE: 1999-10-28
; NUMBER OF SEQ ID NOS: 2626
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 409
; LENGTH: 25821
; TYPE: DNA
; ORGANISM: Drosophila
US-60-161-932-409

```

Query Match	5.2%;	Score 91.8;	DB 88;	Length 25821;
Best Local Similarity	54.3%;	Pred. No. 4.2e-07;		
Matches 208;	Conservative 0;	Mismatches 172;	Indels 3;	Gaps 1;
Qy	163	ttttatctattttttttaaagaaacccagcagataaactctgtcagttattatca	222	
Db	94	tattattattattattattattattattattattattattattattattattattatt	153	
Qy	223	agtcgcatattttcattatcttactcaaatatttgaaattgcatttccaatc	282	
Db	154	attattattattattattattattattattattattattattattattattattatt	213	
Qy	283	taaaatttcaactttttttaacaactttttaaanaattcccttattattatgatttt	342	





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```
RESULT 14
US-09-404-520-13069/c
; Sequence 13069, Application US/09404520
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Ghodessi, Azila
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: McIninch, James
; APPLICANT: Timberlake, William E.
; APPLICANT: Yu, Jaehyuk
; TITLE OF INVENTION: Emericella nidulans Genome Sequence and Uses Thereof
; FILE REFERENCE: 38-10(15498)A
; CURRENT APPLICATION NUMBER: US/09/404,520
; CURRENT FILING DATE: 1999-09-23
; NUMBER OF SEQ ID NOS: 44345
; SEQ ID NO 13069
; LENGTH: 948
; TYPE: DNA
; ORGANISM: Aspergillus nidulans
; FEATURE:
; OTHER INFORMATION: unsure at all n locations
US-09-404-520-13069
```

```
Query Match          5.1%; Score 90.8; DB 52; Length 948;
Best Local Similarity 53.0%; Pred. No. 3.1e-07;
Matches 188; Conservative 0; Mismatches 167; Indels 0; Gaps 0;
```

```
QY 155 attaaagtttactctatctttttttaaaccgaatccaccagcataatcttgcagt 214
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 730 ATTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTT 671

QY 215 attatcaagtcgltatcttttctactactaaatattatgaattgacatt 274
    |||| | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 670 TTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTT 611

QY 275 ccaatcctaaatttaccattttttaaacaattttaaaatttactcttaattat 334
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 610 TTAATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTAT 551

QY 335 agtatcttttttttttttagctaaatataaattctctgaatttattgaatttt 394
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 550 ATATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTAT 491

QY 395 ataattagctactaaataacaaatatttttccctactcagaataaattcatttta 454
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 490 TTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTT 431

QY 455 aaaaaattgattttatagtaatttgltgataattgaattgaattgatt 509
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 430 TTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTATTTTAT 376
```

```
RESULT 15
US-60-146-055-46
; Sequence 46, Application US/60146055
; GENERAL INFORMATION:
; APPLICANT: Bonazzi, Vivien
; TITLE OF INVENTION: ISOLATED SECRETED PROTEINS, NUCLEIC ACID
; TITLE OF INVENTION: MOLECULES ENCODING SECRETED PROTEINS, AND USES THEREOF
; FILE REFERENCE: C1000067
; CURRENT APPLICATION NUMBER: US/60/146,055
```

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; CURRENT FILING DATE: 1999-07-28
; NUMBER OF SEQ ID NOS: 938
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 46
; LENGTH: 1074
; TYPE: DNA
; ORGANISM: Drosophila
US-60-146-055-46
```

```
Query Match          5.0%; Score 89.8; DB 81; Length 1074;
Best Local Similarity 49.9%; Pred. No. 4.8e-07;
Matches 226; Conservative 0; Mismatches 227; Indels 0; Gaps 0;
```

```
QY 49 ttaaatcacattttcttctatcttgcgcgaattttaagaattagtaacataaagta 108
    || ||| ||| | | | | | | | | | | | | | | | | | | | | | | |
DB 460 tttaaagtaatccacgcgcgcgcgaattatataatataatatttttatataatata 519

QY 109 gaacaaagttaattgtaacatttagcgcaatccttaagaanaaattaaagtttat 168
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 520 tacatgtactataatattatattatattatattatattatattatattatattat 579

QY 169 ctatttttttaacgaatccaccagcataaatttgcagatttttcaagtctg 228
    ||| ||| | | | | | | | | | | | | | | | | | | | | | | |
DB 580 ttatattatattatattatattatattatattatattatattatattatattat 639

QY 229 tatttttctattttctactaaatatttgaatttgcatttccataactaaat 288
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 640 ttattattattattatattatattatattatattatattatattatattatattat 699

QY 289 ttacattttttataacaattttaaaatttactctttaaattatagtaatttttatt 348
    || | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 700 ttataattattattatattatattatattatattatattatattatattatattat 759

QY 349 ttttagctaaattaaatatacttgaattttatgaattttataatagtaacta 408
    || ||| | | | | | | | | | | | | | | | | | | | | | | | | |
DB 760 ttacattatattatattatattatattatattatattatattatattatattatatt 819

QY 409 aaaaatacaatatttttccctactcagaanaattcattttttaaaaaaattgatt 468
    | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 820 ttatattattattatattatattatattatattatattatattatattatattat 879

QY 469 ttatagtaatttggtgataattgaattaa 501
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | |
DB 880 ttatattattattatattatattatattat 912
```

Search completed: November 20, 1999, 00:02:14  
Job time: 3343 sec